



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/050,359B TIME: 17:37:41,

Input Set : A:\SequenceListing.txt

Output Set: N:\CRF3\05292002\I050359B.raw

ENTERE

SEQUENCE LISTING

(1) GENERAL INFORMATION:

```
(i) APPLICANT: FOWLKES, Dana M
      6
      7
                             KAY, Brian K
      8
                             FRELINGER, Jeffrey A
      9
                             HYDE-DERUYSCHER, Robin P
            (ii) TITLE OF INVENTION: IDENTIFICATION OF DRUGS USING
     11
                                      COMPLEMENTARY COMBINATORIAL LIBRARIES
     12
           (iii) NUMBER OF SEQUENCES: 178
     14
            (iv) CORRESPONDENCE ADDRESS:
     16
     17
                  (A) ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
     18
                  (B) STREET: 624 Ninth Street N.W., Suite 300
     19
                  (C) CITY: Washington
     20
                  (D) STATE: D.C.
                  (E) COUNTRY: U.S.A.
     21
                  (F) ZIP: 20001
     22
             (V) COMPUTER READABLE FORM:
     24
                  (A) MEDIUM TYPE: Floppy disk
     25
     26
                  (B) COMPUTER: IBM PC compatible
     27
                  (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     28
                  (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
     30
            (vi) CURRENT APPLICATION DATA:
C--> 31
                  (A) APPLICATION NUMBER: US/09/050,359B
C--> 32
                  (B) FILING DATE: 31-Mar-1998
     33
                  (C) CLASSIFICATION:
     39
           (vii) PRIOR APPLICATION DATA:
                  (A) APPLICATION NUMBER: PCT/US97/19638
     36
     37
                  (B) FILING DATE: 31-OCT-1997
     40
                  (A) APPLICATION NUMBER: US 08/740,671
     41
                  (B) FILING DATE: 31-OCT-1996
     43
          (viii) ATTORNEY/AGENT INFORMATION:
     44
                  (A) NAME: COOPER, Iver P
     45
                  (B) REGISTRATION NUMBER: 28,005
     46
                  (C) REFERENCE/DOCKET NUMBER: FOWLKES=4B
     48
            (ix) TELECOMMUNICATION INFORMATION:
     49
                  (A) TELEPHONE: (202) 628-5197
     50
                  (B) TELEFAX: (202) 737-3528
     53
       (2) INFORMATION FOR SEQ ID NO: 1:
     55
             (i) SEQUENCE CHARACTERISTICS:
     56
                  (A) LENGTH: 33 base pairs
     57
                  (B) TYPE: nucleic acid
     58
                  (C) STRANDEDNESS: single
     59
                  (D) TOPOLOGY: linear
```

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61
       (ii) MOLECULE TYPE: cDNA
63
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
65 GANNNKNNKN NKNNKTGGNN KNNKNNKNNK NNK
67 (2) INFORMATION FOR SEQ ID NO: 2:
        (i) SEQUENCE CHARACTERISTICS:
70
             (A) LENGTH: 33 base pairs
71
             (B) TYPE: nucleic acid
72
             (C) STRANDEDNESS: single
73
             (D) TOPOLOGY: linear
75
       (ii) MOLECULE TYPE: cDNA
77
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
79 NNKGANNNKN NKNNKTGGNN KNNKNNKNNK NNK
                                                                            33
81 (2) INFORMATION FOR SEQ ID NO: 3:
        (i) SEQUENCE CHARACTERISTICS:
83
84
             (A) LENGTH: 33 base pairs
             (B) TYPE: nucleic acid
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86
             (C) STRANDEDNESS: single
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             (D) TOPOLOGY: linear
89
       (ii) MOLECULE TYPE: cDNA
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                   33
93 NNKNNKGANN NKNNKTGGNN KNNKNNKNNK NNK
95 (2) INFORMATION FOR SEQ ID NO: 4:
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        (i) SEQUENCE CHARACTERISTICS:
98
             (A) LENGTH: 33 base pairs
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             (B) TYPE: nucleic acid
100
              (C) STRANDEDNESS: single
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              (D) TOPOLOGY: linear
        (ii) MOLECULE TYPE: cDNA
102
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
104
106 NNKNNKNNKG ANNNKTGGNN KNNKNNKNNK NNK
                                                                    33
108 (2) INFORMATION FOR SEQ ID NO: 5:
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         (i) SEQUENCE CHARACTERISTICS:
111
            (A) LENGTH: 33 base pairs
              (B) TYPE: nucleic acid
112
113
              (C) STRANDEDNESS: single
114
              (D) TOPOLOGY: linear
116
        (ii) MOLECULE TYPE: cDNA
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
118
                                                                    33
120 NNKNNKNNKN NKGANTGGNN KNNKNNKNNK NNK
122 (2) INFORMATION FOR SEQ ID NO: 6:
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         (i) SEQUENCE CHARACTERISTICS:
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              (A) LENGTH: 33 base pairs
126
              (B) TYPE: nucleic acid
127
              (C) STRANDEDNESS: single
128
              (D) TOPOLOGY: linear
130
        (ii) MOLECULE TYPE: cDNA
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        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
134 NNKNNKNNKN NKNNKTGGGA NNNKNNKNNK NNK
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136 (2) INFORMATION FOR SEQ ID NO: 7:

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138
               (i) SEQUENCE CHARACTERISTICS:
     139
                    (A) LENGTH: 33 base pairs
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                    (B) TYPE: nucleic acid
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                    (C) STRANDEDNESS: single
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                    (D) TOPOLOGY: linear
     143
             (ii) MOLECULE TYPE: cDNA
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             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
     147 NNKNNKNNKN NKNNKTGGNN KGANNNKNNK NNK
                                                                          33
     150 (2) INFORMATION FOR SEQ ID NO: 8:
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              (i) SEQUENCE CHARACTERISTICS:
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                    (A) LENGTH: 33 base pairs
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                    (B) TYPE: nucleic acid
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                    (C) STRANDEDNESS: single
     156
                    (D) TOPOLOGY: linear
     157
             (ii) MOLECULE TYPE: cDNA
     159
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
     161 NNKNNKNNKN NKNNKTGGNN KNNKGANNNK NNK
                                                                          33
     163 (2) INFORMATION FOR SEQ ID NO: 9:
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              (i) SEQUENCE CHARACTERISTICS:
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                   (A) LENGTH: 33 base pairs
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                   (B) TYPE: nucleic acid
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                   (C) STRANDEDNESS: single
     169
                   (D) TOPOLOGY: linear
     171
             (ii) MOLECULE TYPE: cDNA
     173
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
     175 NNKNNKNNKN NKNNKTGGNN KNNKNNKGAN NNK
                                                                         33
     177 (2) INFORMATION FOR SEQ ID NO: 10:
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              (i) SEQUENCE CHARACTERISTICS:
     180
                   (A) LENGTH: 33 base pairs
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                   (B) TYPE: nucleic acid
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                   (C) STRANDEDNESS: single
     183
                   (D) TOPOLOGY: linear
     184
             (ii) MOLECULE TYPE: cDNA
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
     188 NNKNNKNNKN NKNNKTGGNN KNNKNNKNNK GAN
                                                                         33
     190 (2) INFORMATION FOR SEQ ID NO: 11:
     192
              (i) SEQUENCE CHARACTERISTICS:
     193
                   (A) LENGTH: 10 amino acids
     194
                   (B) TYPE: amino acid
     195
                   (C) STRANDEDNESS: single
     196
                   (D) TOPOLOGY: linear
             (ii) MOLECULE TYPE: peptide
     197
     199
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
W--> 201
              Xaa Xaa Xaa Xaa Xyr Xaa Xaa Xaa Xaa
     202
              1
                               5
     204 (2) INFORMATION FOR SEQ ID NO: 12:
     206
              (i) SEQUENCE CHARACTERISTICS:
     207
                   (A) LENGTH: 10 amino acids
     208
                   (B) TYPE: amino acid
```

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Input Set : A:\SequenceListing.txt Output Set: N:\CRF3\05292002\I050359B.raw 209 (C) STRANDEDNESS: single 210 (D) TOPOLOGY: linear 211 (ii) MOLECULE TYPE: peptide (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12: 213 Xaa Xaa Xaa Xaa Tyr Leu Xaa Xaa Xaa W--> 215216 5 (2) INFORMATION FOR SEQ ID NO: 13: 218 220 (i) SEQUENCE CHARACTERISTICS: 221 (A) LENGTH: 10 amino acids 222 (B) TYPE: amino acid 223 (C) STRANDEDNESS: single (D) TOPOLOGY: linear 224 226 (ii) MOLECULE TYPE: peptide 228 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13: W--> 230Xaa Xaa Xaa Pro Pro Xaa Xaa Pro Xaa Xaa 231 233 (2) INFORMATION FOR SEQ ID NO: 14: 235 (i) SEQUENCE CHARACTERISTICS: 236 (A) LENGTH: 14 amino acids 237 (B) TYPE: amino acid 238 (C) STRANDEDNESS: single 239 (D) TOPOLOGY: linear 241 (ii) MOLECULE TYPE: peptide 243 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14: W--> 245 Xaa Xaa Xaa Xaa Xaa Pro Xaa Pro Pro Xaa Pro Xaa Xaa 246 5 10 248 (2) INFORMATION FOR SEQ ID NO: 15: 250 (i) SEQUENCE CHARACTERISTICS: 251 (A) LENGTH: 27 base pairs 252 (B) TYPE: nucleic acid 253 (C) STRANDEDNESS: single 254 (D) TOPOLOGY: linear 256 (ii) MOLECULE TYPE: cDNA 258 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15: 27 260 CTGTGCGGAT CCATGGATCG CAAGACG 262 (2) INFORMATION FOR SEQ ID NO: 16: 264 (i) SEQUENCE CHARACTERISTICS: 265 (A) LENGTH: 27 base pairs 266 (B) TYPE: nucleic acid 267 (C) STRANDEDNESS: single 268 (D) TOPOLOGY: linear 270 (ii) MOLECULE TYPE: cDNA 272 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16: 27 274 CTGTGCGAAT TCCTAGCCGC ACTTTTG 276 (2) INFORMATION FOR SEQ ID NO: 17: 278 (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 63 base pairs

(C) STRANDEDNESS: single

(B) TYPE: nucleic acid

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Input Set : A:\SequenceListing.txt Output Set: N:\CRF3\05292002\I050359B.raw 282 (D) TOPOLOGY: linear 284 (ii) MOLECULE TYPE: cDNA (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17: 286 288 GACTGTGCCT CGAGKNNKNN KNNKNNKNNK YYYNNKNNKN NKNNKNNKTC TAGACGTGTC 60 290 AGT 63 292 (2) INFORMATION FOR SEQ ID NO: 18: (i) SEQUENCE CHARACTERISTICS: 295 (A) LENGTH: 15 base pairs 296 (B) TYPE: nucleic acid 297 (C) STRANDEDNESS: single 298 (D) TOPOLOGY: linear 300 (ii) MOLECULE TYPE: cDNA 302 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18: 304 ACTGACACGT CTAGA 15 306 (2) INFORMATION FOR SEQ ID NO: 19: 308 (i) SEQUENCE CHARACTERISTICS: 309 (A) LENGTH: 22 amino acids 310 (B) TYPE: amino acid 311 (C) STRANDEDNESS: single 312 (D) TOPOLOGY: linear 314 (ii) MOLECULE TYPE: peptide 316 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: W--> 318Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Glu His Val Cys Ser 319 321 Trp Gly Trp Gly Arg Cys 322 20 324 (2) INFORMATION FOR SEQ ID NO: 20: (i) SEQUENCE CHARACTERISTICS: 327 (A) LENGTH: 11 amino acids 328 (B) TYPE: amino acid 329 (C) STRANDEDNESS: single 330 (D) TOPOLOGY: linear 332 (ii) MOLECULE TYPE: peptide 334 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: 336 Pro Thr Ser Asp Leu Trp Arg Asn Leu Gly Gly 337 5 10 339 (2) INFORMATION FOR SEQ ID NO: 21: 341 (i) SEQUENCE CHARACTERISTICS: 342 (A) LENGTH: 11 amino acids 343 (B) TYPE: amino acid 344 (C) STRANDEDNESS: single 345 (D) TOPOLOGY: linear 347 (ii) MOLECULE TYPE: peptide 349 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21: 351 Trp Gly Glu Thr Met Trp Asp Asn Arg Lys Val 352 1 5 10 354 (2) INFORMATION FOR SEQ ID NO: 22: 356 (i) SEQUENCE CHARACTERISTICS: 357 (A) LENGTH: 11 amino acids

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